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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/964,824C

DATE: 06/03/2003 P.6
TIME: 13:40:14

Input Set : A:\Avalon 73.txt
Output Set: N:\CRF4\06032003\I964824C.raw

4 <110> APPLICANT: Horrigan, Stephen
 6 <120> TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
 7 Signature Gene Sets
 9 <130> FILE REFERENCE: 689290-73
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/964,824C
 C--> 11 <141> CURRENT FILING DATE: 2001-09-27
 11 <150> PRIOR APPLICATION NUMBER: US/60/236,033
 12 <151> PRIOR FILING DATE: 2000-09-28
 14 <150> PRIOR APPLICATION NUMBER: US/60/236,032
 15 <151> PRIOR FILING DATE: 2000-09-28
 17 <150> PRIOR APPLICATION NUMBER: US/60/236,028
 18 <151> PRIOR FILING DATE: 2000-09-28
 20 <160> NUMBER OF SEQ ID NOS: 583
 22 <170> SOFTWARE: PatentIn version 3.0
 24 <210> SEQ ID NO: 1
 25 <211> LENGTH: 521
 26 <212> TYPE: DNA
 27 <213> ORGANISM: Homo sapiens
 29 <220> FEATURE:
 30 <221> NAME/KEY: misc_feature
 31 <222> LOCATION: (1)...(521)
 32 <223> OTHER INFORMATION: n=a,t,g or c
 35 <400> SEQUENCE: 1
 36 gtaatatggaa attagaaaaca atttggcttt ttagagctga aactagaaaac aacacatcca 60
 38 ggaacagtag acttcttattt ctttcaatcc ctaatgtcct agtgagtatg taccctatgg 120
 40 agaaggcaga aatgacgtgg accaggactc cttacatggaa gagtgtttt aaggcagttt 180
 42 taaaaaagcc cattttgtga aagaaaccag aaggctcgta attgctgtct gcactgtgg 240
 W--> 44 ttctcctgggg ggttggggag gggagtggat taaataaaaaa gtttagaagg ccatagnata 300
 46 aatatcgaaa tagtatgaat tttatatat atctttaaag gggtaggca atgatgaaaa 360
 48 gatatgactg ctttccttcc atttctcatt aaattaaaaat tcccacaaaaa gtgcacggca 420
 50 tcttttgaa acactgctaa ttttaaagg ttggaaaggtt tatcttcata gccacaaatct 480
 52 ttgcnaaagg cttggtaccg gnaacaaggc tccagtctgc c 521
 55 <210> SEQ ID NO: 2
 56 <211> LENGTH: 481
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Homo sapiens
 60 <400> SEQUENCE: 2
 61 ataaaatggtt ttttttaac ataagtaaat ttacaaatca aatgaaaaat gaaaaataca 60
 63 aaagttcatg aatgaaataa aaaagacact cttcaaatat taaaacctat ggaaagaaaa 120
 65 taagtaatta atgaatgtat tttttgtttc caaatacaat gaagtgattt tttatttagag 180
 67 tccttggaa tcatctaagt tacaatacag aagagaatta aataaaatgt atatgatttt 240
 69 gtaatttagac actctatata tcacagttt ttgttaacct gggcatggaa cgtccctata 300
 71 gcatatattt aaaaccatta atttttttt aaaaaatttg agacatggtt tgttcttgg 360

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73 ctctaaatata tttttcccca tttcccttga atgttctcta ttggccatct tctggAACAT 420
 75 taaaaaaaaaa tcttggAAACA aattcttcttG caatgatacG tatcacataa acttgatATG 480
 77 C 481

80 <210> SEQ ID NO: 3
 81 <211> LENGTH: 357
 82 <212> TYPE: DNA
 83 <213> ORGANISM: Homo sapiens
 85 <400> SEQUENCE: 3

86 gagcgggtgga gggcgtcaCT gggtttgcgc gtctggcaag cgattcagCT gtctgctccc 60
 88 tagcagccgg ccttcgggtc gggcgtcttc cccggtact gcccgttca ttcttccgg 120
 90 gtggccacga gtcgggttgc acttctgtga tccatcctca tcttctaaAG atgcatcctg 180
 92 acttatctcc acacttgcac actgaagaAT gcaacgttCT gattaacttG cttaaggAA 240
 94 gtcacaaaaaa tcacaacattt ctgaaatttt ttgggtatttG taatgatgtt gatcgggggt 300
 96 ggagagaatg cctctagAGT gatgtacata gagaacagGA gcccggcAG ggggcat 357

99 <210> SEQ ID NO: 4
 100 <211> LENGTH: 1086
 101 <212> TYPE: DNA
 102 <213> ORGANISM: Homo sapiens
 104 <400> SEQUENCE: 4

105 cgcagccccc cgcccccccc ctcagcggcc ggccccggGA tgacggcggc ccaggccggc 60
 107 ggtgaggagg cgccaccagg cgtgcgggtc gtcaagggtgg ttctgggtgg cgacggcggc 120
 109 tgcgggaaAGA cgtcgctgt gatggtcttc gccgatgggg cttccccGA gagctacacc 180
 111 cccacgggtt ttgagcggta catggtcaac ctgcaagtGA aaggcaAAAC tgcacccctc 240
 113 cacatctggg acacaggcagg gcaagatgac tatgaccggc tgcggccctt gttctaccct 300
 115 gacggcagcg tcctgtgt gtcaccaggc cgaacagCT ttctgcaAGA aggtaccatC catcgctgt 360
 117 tttaaccggT ggtacccAGA agtgaatcat ttctgcaAGA aggtaccatC catcgctgt 420
 119 ggctgcaAGA ctgacccgtcg caaggacaaa tcactggtGA acaagctccG aagaaacggA 480
 121 ttggagccTg tgacccatca cagggggccAG gagatggcGA ggtccgtggg cgccggggcc 540
 123 tacctcgagt gctcggtcg gctccatgac aacgtccacg ccgtcttcca ggaggccggc 600
 125 gaggtggccc tcagcggcccg cggtcgcaac ttctggccggc ggattaccca gggctttgc 660
 127 gtggtgacG gagcggctcg gggcgtccca ggcacgggg aaggggcagg ggcgtgacct 720
 129 gctgctgagC tggctgggt ggaccggcgtc cctaggctgt gaccggcggaa ctccactgca 780
 131 acagacgggc gccaccaaaAG ccaggccctg aggctggGA gtcctggact gagaaggGGG 840
 133 gttccctggc ccacccgtcg tggtagggc tcgtccctgc gtgcccggAG atcaactcgct 900
 135 aaccctatg cccgggtcccG gaccgacatc ctggagccgc ctgtgcAGCC tgatgcccc 960
 137 tcgtggctgc tcccagggtc gcacccgtca ggacctaATG ttcttaggtc cctctggcca 1020
 139 gaacccacac ccggccccctt cccacccgtc atactggtaA ctgtacaAGA aaaaacgaca 1080
 141 tcactt 1086

144 <210> SEQ ID NO: 5
 145 <211> LENGTH: 486
 146 <212> TYPE: DNA
 147 <213> ORGANISM: Homo sapiens
 149 <220> FEATURE:
 150 <221> NAME/KEY: misc_feature
 151 <222> LOCATION: (1)...(486)
 152 <223> OTHER INFORMATION: n=a,t,g or c
 155 <400> SEQUENCE: 5

156 tagcaccatG atccctcgccG tggagctgtG tgaggAGATC gtggctatG ggatggtcAG 60
 W--> 158 cgacantanc tgcagggAGA agggccacCC ctcagtgcct taccactact ttgagaaggG 120

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252	cgtgcttcat	ccctttattt	aagaacagaa	aattatgact	actctacaag	gtggataata	180
254	ttttggtaacc	tgtgctggc	cacagccctg	ttcctcaaag	ctgaattttag	agatttctct	240
256	ttgacttcca	agacccatgc	gttataaggc	acccctgaaat	aaatttttg	tgccctggaaa	300
W--> 258	tgcagggagg	gcaatagctt	tgtaaatgg	tttacat	tttctgtaa	tttttcttagg	360
260	gtccttagtgc	ttccgaatca	tttaatgc	ttgtcgata	tccttttaca	tttcaattgc	420
262	aatccatgaa	attacattt	gaagattt	agtacttaac	ggtagtctc	ccatgaattt	480
265	<210>	SEQ ID NO:	8				
266	<211>	LENGTH:	416				
267	<212>	TYPE:	DNA				
268	<213>	ORGANISM:	Homo sapiens				
270	<220>	FEATURE:					
271	<221>	NAME/KEY:	misc_feature				
273	<222>	LOCATION:	(1)...(416)				
274	<223>	OTHER INFORMATION:	n=a,t,g or c				
277	<400>	SEQUENCE:	8				
W--> 278	atttcagang	aagtttatta	agaggttta	ggctttaagc	atatgtgaaa	agcaaaaatt	60
280	acattttaaa	gtatataatt	tgcattttcc	accttctcaa	tgccaatgaa	atattctagg	120
282	agactctata	agataaccaa	ttgattttct	actactccc	aattttact	ttgttaattt	180
284	aagaggaata	ggcaaata	gctgctgtgg	ttctggttct	ccctgcagga	tgaagggggc	240
286	ctgcaaaatg	tctccactt	ccattctagg	tcattcagca	aggtgccttc	ctctggatgc	300
288	actgtctgt	tacttttgc	atgttgcatc	acataatgga	ttctggccca	ccttacacca	360
290	ttttgactgt	cagtaaaaga	atggatgtgt	ggcccat	ttcn	tttattt aatagc	416
293	<210>	SEQ ID NO:	9				
294	<211>	LENGTH:	371				
295	<212>	TYPE:	DNA				
296	<213>	ORGANISM:	Homo sapiens				
298	<400>	SEQUENCE:	9				
299	tttgacacgt	gaagggttat	ttatggttat	gatgaccctg	tcctgcaacg	aggactggc	60
301	agccactact	gaggaggagg	gtcccatctc	tctcctgtcg	gcttcaccg	aggcacagc	120
303	cagacgtgg	gcaaagggt	tccctgtct	accaggccat	tcctggcct	gccgcctagg	180
305	gctcacagg	gcccaggagt	ccccagctca	cagccagg	catcaggcca	ggcgcgtcg	240
307	gtgcacaccg	cacctgtgg	ggacctgggt	acactcagga	gaccaagagc	actggcggt	300
309	caggatgg	ggcgttcagc	tcctacgggg	tgggagaag	tctgtagccg	agagcccagc	360
311	ccccctcctgc	c					371
314	<210>	SEQ ID NO:	10				
315	<211>	LENGTH:	419				
316	<212>	TYPE:	DNA				
317	<213>	ORGANISM:	Homo sapiens				
319	<400>	SEQUENCE:	10				
320	aagtattctg	tcccttaat	agctttgttt	tagggtaac	tcccctcgcc	ttgtggggag	60
322	gcttaggacg	ggcggtgtca	atcctcgaag	gggagtctca	gcgaccatgg	ggacaccat	120
324	ccacatgcag	gcccgtatgt	gggcctcg	agccctctc	ttgggtgtcc	ggggccctcg	180
326	ttgcccctca	gtgcctgt	gcaagg	ttgcagggcc	ggctgttacc	gctgaggctc	240
328	ttggatattt	gagatggagg	attctggctc	aggagtctca	gggagtgc	ctgaaggag	300
330	gtgatattt	gggcccgtc	aatgaggatc	caggtgacag	tctcggagca	gggcgggt	360
332	ctgagagagc	cctgataggt	gatgaaaccg	aagattcagg	gaacaggagc	tccaggctc	419
335	<210>	SEQ ID NO:	11				
336	<211>	LENGTH:	270				
337	<212>	TYPE:	DNA				

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338 <213> ORGANISM: Homo sapiens
 340 <400> SEQUENCE: 11
 341 tacagggcaa cccaccccta ggcaaaggct cggcctctcc cacctcccc acgtcatcac 60
 343 tgagctgcgg cacgcagagg tgccagccaa ttccgaggag aattgggtcc aatagaaata 120
 345 tttacaaata accagggggc aggtgtcccg tgatccggaa tcgtgagggg acttagtacc 180
 347 agggggccct tggctcccaa cagccccagg ccctggggcg gacttggcac aggacccaag 240
 349 agggaaactgg ggcattgggg ggccggcaga 270
 352 <210> SEQ ID NO: 12
 353 <211> LENGTH: 255
 354 <212> TYPE: DNA
 355 <213> ORGANISM: Homo sapiens
 357 <400> SEQUENCE: 12
 358 tttagtttag caccatttat taagtgatct cagctgtgt ttagctgct gcgtgtcacc 60
 360 gtgttcttaa aacataaaat gctcttccga ttcccttctgt ccaggacaga aggttcttcc 120
 362 aggttagcact ccaacagaac aagagactcc gatgacgcca gcttcaatga tggtgccatc 180
 364 cagagagggg gaggttcatg gcacattcaa ccgcggcttc cagaggtttt gaaaaaggag 240
 366 ctttggggg cccag 255
 369 <210> SEQ ID NO: 13
 370 <211> LENGTH: 358
 371 <212> TYPE: DNA
 372 <213> ORGANISM: Homo sapiens
 374 <400> SEQUENCE: 13
 375 caggttgaat aaaatttat tgataatgct ttatattaat attctctttt gcatttaat 60
 377 attatatgaa tactacaagg atccaacaag aaataacctt cataaatttag cataattttat 120
 379 agcagggaaac ccaaataaaac taaacttggc tgcctaaat aatttgcata aagggggata 180
 381 tgctctttgt aagtatcatg ctgataaaac cataaaaatt ctttagagg aatgaggatt 240
 383 aaaatgaaaat ttctttatga cacggaaaaa aataataatt tgcataaaag tggaaaattt 300
 385 taaaagcaaa cattatacac ataaccagca caattatttc catcttaaaa cattggtt 358
 388 <210> SEQ ID NO: 14
 389 <211> LENGTH: 266
 390 <212> TYPE: DNA
 391 <213> ORGANISM: Homo sapiens
 393 <400> SEQUENCE: 14
 394 atggctaattt gtgacacact ttatattttt aaaaacacgc cttccacaca tagtgcgtga 60
 396 ggcatgtgca cattttccta gaaggacatg aatagtgtat tggaggtacg gtggaggtca 120
 398 ggcatctaca gggtcattcg aggaggaaca gattcaagct ttcggacgat cagtgtttt 180
 400 taaatagcag catcatcaga tctaagacaa cattggacct ggcagggcct tttctttggg 240
 402 tggcattaat tactccagat tcagac 266
 405 <210> SEQ ID NO: 15
 406 <211> LENGTH: 287
 407 <212> TYPE: DNA
 408 <213> ORGANISM: Homo sapiens
 410 <400> SEQUENCE: 15
 411 aacgtaaaca caaagtctca ttatattttt tctgaaggcac acaggagctc actcagcaca 60
 413 ataacagtaa gcaatcata caaatattga gaaaaatgt tcctatgtat acatacatgt 120
 415 atattcttaa gagtagcgtt caggagttt acaacaaatg taaagtgggt ttctctaaag 180
 417 aatgtttctt gacaggctt tgggttggaa atggacaggt aaatcactgt cacataacag 240
 419 gtaagctaag aataacttct gttacccaag tcatttgaac cctgtgg 287
 422 <210> SEQ ID NO: 16

RAW SEQUENCE LISTING ERROR SUMMARY
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 297,485,502
Seq#:5; N Pos. 66,69,389,483
Seq#:7; N Pos. 331
Seq#:8; N Pos. 9,404
Seq#:16; N Pos. 113,115,248,252
Seq#:17; N Pos. 15,386
Seq#:21; N Pos. 90,258
Seq#:22; N Pos. 330,361,419
Seq#:23; N Pos. 371,416
Seq#:24; N Pos. 217,232,315,333
Seq#:27; N Pos. 26,199
Seq#:28; N Pos. 145,222,228
Seq#:51; N Pos. 92,118,152,153,235,237,238,239,244,245,248,263,270,284
Seq#:52; N Pos. 12,352,359,400,408
Seq#:54; N Pos. 346,368
Seq#:55; N Pos. 180
Seq#:56; N Pos. 27
Seq#:57; N Pos. 432,478,495
Seq#:59; N Pos. 481
Seq#:63; N Pos. 104,120,188
Seq#:64; N Pos. 44,146,265,285
Seq#:65; N Pos. 11,130,133,190,313
Seq#:66; N Pos. 375,417
Seq#:68; N Pos. 146
Seq#:69; N Pos. 187,223
Seq#:70; N Pos. 3
Seq#:71; N Pos. 158,161,274,393
Seq#:77; N Pos. 8,10,11,21,28,37,54,55,117,126,130,133,152,154,155,158,159
Seq#:77; N Pos. 160,161,164,179,180,187,193,335,417,461,503,526
Seq#:78; N Pos. 398
Seq#:79; N Pos. 519,535
Seq#:80; N Pos. 575
Seq#:81; N Pos. 167
Seq#:90; N Pos. 274,366,371,458,463,496,504,510,519
Seq#:91; N Pos. 12,89,109,191,468
Seq#:93; N Pos. 2,3,35,158,199,238,271,421,459,524,540
Seq#:107; N Pos. 503,512
Seq#:108; N Pos. 343
Seq#:118; N Pos. 59,315,325,330,343,355,363,369,394
Seq#:123; N Pos. 410
Seq#:124; N Pos. 332,463,492,510,525,541,546,551,589,606,622
Seq#:125; N Pos. 515
Seq#:126; N Pos. 3,4,5,12,174,390,401,429,483,524,530,532,533,534,535,536
Seq#:126; N Pos. 537,538,539,540,541,542,543,544,545,546,547,548,549,550

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Seq#:126; N Pos. 551,552,553,554,556,557,560,561,568,569,570,572,573,574
Seq#:126; N Pos. 575,576,577,578,579,580,581,583,584,585,586,587,588,589
Seq#:126; N Pos. 591,593,594,595,597,598,599,600,602,603,604,606,607,609
Seq#:126; N Pos. 610,611,612,613,614,615,616,619,620,621,622,624,625,626
Seq#:126; N Pos. 627,631,632,633,634,635,636,638,639,641,643,644,646,648
Seq#:126; N Pos. 649,651,652,653,654,655,656,659,662,663,665,667,668,669
Seq#:126; N Pos. 671,674,675,677,678,679,680,681,682,683,684,686,688,690

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L:11 M:270 C: Current Application Number differs, Replaced Current Application No
 L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
 L:44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:240
 M:341 Repeated in SeqNo=1
 L:158 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:60
 M:341 Repeated in SeqNo=5
 L:258 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:300
 L:278 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
 M:341 Repeated in SeqNo=8
 L:436 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:60
 M:341 Repeated in SeqNo=16
 L:458 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:0
 M:341 Repeated in SeqNo=17
 L:539 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:60
 M:341 Repeated in SeqNo=21
 L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:300
 M:341 Repeated in SeqNo=22
 L:605 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:360
 L:631 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:180
 M:341 Repeated in SeqNo=24
 L:695 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
 M:341 Repeated in SeqNo=27
 L:729 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28 after pos.:120
 M:341 Repeated in SeqNo=28
 L:1209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:60
 M:341 Repeated in SeqNo=51
 L:1231 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52 after pos.:0
 M:341 Repeated in SeqNo=52
 L:1288 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54 after pos.:300
 M:341 Repeated in SeqNo=54
 L:1310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:55 after pos.:120
 L:1330 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56 after pos.:0
 L:1374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:57 after pos.:420
 M:341 Repeated in SeqNo=57
 L:1431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59 after pos.:480
 L:1504 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:63 after pos.:60
 M:341 Repeated in SeqNo=63
 L:1526 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:64 after pos.:0
 M:341 Repeated in SeqNo=64
 L:1552 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:65 after pos.:0
 M:341 Repeated in SeqNo=65
 L:1590 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:66 after pos.:360
 L:1631 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:68 after pos.:120
 L:1657 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:69 after pos.:180
 L:1675 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:70 after pos.:0
 L:1690 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
 L:1699 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:71 after pos.:120
 M:341 Repeated in SeqNo=71

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L:1820 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:77 after pos.:0
M:341 Repeated in SeqNo=77
L:1866 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:78 after pos.:360
L:1900 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:79 after pos.:480
L:1936 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:80 after pos.:540



Creation date: 11-28-2003

Indexing Officer: MALI3 - MOHAMED ALI

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Dossier: 09964824

Legal Date: 06-10-2003

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